C/Æ   CSC
CSC

CRF Errors Corrected by the STIC Systems Branch.

Serial Number: 09/925,055D

ENTERED

CRF Processing Date: 3/17/2003
Edited by: ______
Verified by: ______ (STIC staff)

- [ ] Changed a file from non-ASCII to ASCII
- [ ] Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- [ ] Edited a format error in the Current Application Data section, specifically: ______
- [ ] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other ______.
- [ ] Added the mandatory heading and subheadings for "Current Application Data".
- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: ______
- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: ______
- [ ] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: ______
- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- [ ] Inserted colons after headings/subheadings. Headings edited included: ______
- [ ] Deleted extra, invalid, headings used by an applicant, specifically: ______
- [x] Deleted: [x] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as ______.
- [ ] Inserted mandatory headings, specifically: ______
- [ ] Corrected an obvious error in the response, specifically: ______
- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.
- [ ] Corrected an error in the Number of Sequences field, specifically: ______
- [ ] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: ______
- [ ] Other: ______

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING DATE: 03/17/2003
PATENT APPLICATION: **US/09/925,055D** TIME: 13:20:10

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\03172003\I925055D.raw**

<1160> APPLICANT: Findsvogel, Wayne R.
Topouzi, Stavroa
<120> TITLE OF INVENTION: SOLUBLE ZCYTOR11 CYTOKINE RECEPTORS
<130> FILE REFERENCE: 50-56
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/925,055D
C--> 14 <141> CURRENT FILING DATE: 2001-08-08
<150> PRIOR APPLICATION NUMBER: US 60/223,827
<151> PRIOR FILING DATE: 2000-08-08
<150> PRIOR APPLICATION NUMBER: US 60/230,876
<151> PRIOR FILING DATE: 2000-12-01
<160> NUMBER OF SEQ ID NOS: 35
<170> SOFTWARE: FastSEQ for Windows Version 3.0
<210> SEQ ID NO: 1
<211> LENGTH: 2851
<212> TYPE: DNA
<213> ORGANISM: Homo sapien
<220> FEATURE:
<221> NAME/KEY: CDS
<222> LOCATION: (548...1755)
<400> SEQUENCE: 1

```
tagagnccaa gngaggnctc tgtgccagcc ccg atg agg acg ctg ctg acc atc        54
                                      Met Arg Thr Leu Leu Thr Ile
                                       1               5

ttg act gtg gga tcc ctg gct gct cac gcc cct gag gac ccc tcg gat       102
Leu Thr Val Gly Ser Leu Ala Ala His Ala Pro Glu Asp Pro Ser Asp
            10                  15                  20

ctg ctc cag cac gtc aaa ttc cag tcc agc aac ttt gaa aac atc ctg       150
Leu Leu Gln His Val Lys Phe Gln Ser Ser Asn Phe Glu Asn Ile Leu
        25                  30                  35

acg tgg gac agc ggc cca gag ggc acc ccg gac acg gtc tac agc atc       198
Thr Trp Asp Ser Gly Pro Glu Gly Thr Pro Asp Thr Val Tyr Ser Ile
    40                  45                  50                  55

gag tat aaa acg tac gga gag agg gac tgg gtg gca aag aag ggc tgt       246
Glu Tyr Lys Thr Tyr Gly Glu Arg Asp Trp Val Ala Lys Lys Gly Cys
                60                  65                  70

cag cgg atc acc cgg aag tcc tgc aac ctg acg gtg gag acg ggc aac       294
Gln Arg Ile Thr Arg Lys Ser Cys Asn Leu Thr Val Glu Thr Gly Asn
            75                  80                  85

ctc acg gag ctc tac tat gcc agg gtc acc gct gtc agt gcg gga ggc       342
Leu Thr Glu Leu Tyr Tyr Ala Arg Val Thr Ala Val Ser Ala Gly Gly
        90                  95                  100

cgg tca gcc acc aag atg act gac agg ttt agc tct ctg cag cac act       390
Arg Ser Ala Thr Lys Met Thr Asp Arg Phe Ser Ser Leu Gln His Thr
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/925,055D

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\03172003\I925055D.raw

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64      125                 110                 115
66  acc ctc aag cca cct gat gtg acc tgt atc tcc aaa gtg aga tcg att      438
67  Thr Leu Lys Pro Pro Asp Val Thr Cys Ile Ser Lys Val Arg Ser Ile
68  120                 125                 130                 135
70  cag atg att gtt cat cct acc ccc acg cca atc cgt gca ggc gat ggg      486
71  Gln Met Ile Val His Pro Thr Pro Thr Pro Ile Arg Ala Gly Asp Gly
72                  140                 145                 150
74  cac cgg cta acc ctg gaa gac atc ttc cat gac ctg ttc tac cac tta      534
75  His Arg Leu Thr Leu Glu Asp Ile Phe His Asp Leu Phe Tyr His Leu
76              155                 160                 165
78  gag ctc cag gtc aac cgc acc tac caa atg cac ctt gga ggg aag cag      582
79  Glu Leu Gln Val Asn Arg Thr Tyr Gln Met His Leu Gly Gly Lys Gln
80          170                 175                 180
82  aga gaa tat gag ttc ttc ggc ctg acc cct gac aca gag ttc ctt ggc      630
83  Arg Glu Tyr Glu Phe Phe Gly Leu Thr Pro Asp Thr Glu Phe Leu Gly
84      185                 190                 195
86  acc atc atg att tgc gtt ccc acc tgg gcc aaq gag agt gcc ccc tac      678
87  Thr Ile Met Ile Cys Val Pro Thr Trp Ala Lys Glu Ser Ala Pro Tyr
88  200                 205                 210                 215
90  atg tgc cga gtg aag aca ctg cca gac cgg aca tgg acc tac tcc ttc      726
91  Met Cys Arg Val Lys Thr Leu Pro Asp Arg Thr Trp Thr Tyr Ser Phe
92                  220                 225                 230
94  tcc gga gcc ttc ctg ttc tcc atg ggc ttc ctc gtc gca gta ctc tgc      774
95  Ser Gly Ala Phe Leu Phe Ser Met Gly Phe Leu Val Ala Val Leu Cys
96              235                 240                 245
98  tac ctg agc tac aga tat gtc acc aaq ccq cct gca cct ccc aac tcc      822
99  Tyr Leu Ser Tyr Arg Tyr Val Thr Lys Pro Pro Ala Pro Pro Asn Ser
100         250                 255                 260
102 ctg aac gtc cag cga gtc ctg act ttc cag ccg ctg cgc ttc atc cag      870
103 Leu Asn Val Gln Arg Val Leu Thr Phe Gln Pro Leu Arg Phe Ile Gln
104     265                 270                 275
106 gag cac gtc ctg atc cct gtc ttt gac ctc agc ggc ccc agc agt ctg      918
107 Glu His Val Leu Ile Pro Val Phe Asp Leu Ser Gly Pro Ser Ser Leu
108 280                 285                 290                 295
110 gcc cag cct gtc cag tac tcc cag atc agg gtg tct gga ccc agg gag      966
111 Ala Gln Pro Val Gln Tyr Ser Gln Ile Arg Val Ser Gly Pro Arg Glu
112                 300                 305                 310
114 ccc gca gga gct cca cag cgg cat agc ctg tcc gag atc acc tac ttg     1014
115 Pro Ala Gly Ala Pro Gln Arg His Ser Leu Ser Glu Ile Thr Tyr Leu
116             315                 320                 325
118 ggg cag cca gac atc tcc atc ctc cag ccc tcc aac gtg cca cct ccc     1062
119 Gly Gln Pro Asp Ile Ser Ile Leu Gln Pro Ser Asn Val Pro Pro Pro
120         330                 335                 340
122 cag atc ctc tcc cca ctg tcc tat gcc cca aac gct gcc cct gag gtc     1110
123 Gln Ile Leu Ser Pro Leu Ser Tyr Ala Pro Asn Ala Ala Pro Glu Val
124     345                 350                 355
126 ggg ccc cca tcc tat gca cct cag gtg acc ccc gaa act caa ttc cca     1158
127 Gly Pro Pro Ser Tyr Ala Pro Gln Val Thr Pro Glu Ala Gln Ile Pro
128 360                 365                 370                 375
```

RAW SEQUENCE LISTING DATE: 03/17/2003
PATENT APPLICATION: US/09/925,055D TIME: 13:20:10

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\03172003\I925055D.raw

```
120  ttc tac gcc cca cag gcc atc tct aag gtc cag cct tcc tcc tat gcc    1206
121  Phe Tyr Ala Pro Gln Ala Ile Ser Lys Val Gln Pro Ser Ser Tyr Ala
122                  380                 385                 390
123  cct caa gcc act ccg gac agc tgg cct ccc tcc tat ggg gta tgc atg    1254
124  Pro Gln Ala Thr Pro Asp Ser Trp Pro Pro Ser Tyr Gly Val Cys Met
125              395                 400                 405
126  gaa ggt tct ggc aaa gac tcc ccc act ggg aca ctt tct agt cct aaa    1302
127  Glu Gly Ser Gly Lys Asp Ser Pro Thr Gly Thr Leu Ser Ser Pro Lys
128          410                 415                 420
129  cac ctt agg cct aaa ggt cag ctt cag aaa gag cca ccg gct gga agc    1350
130  His Leu Arg Pro Lys Gly Gln Leu Gln Lys Glu Pro Pro Ala Gly Ser
131      425                 430                 435
132  tgc atg tta ggt ggc ctt tct ctg cag gag gtg acc tcc ttg gct atg    1398
133  Cys Met Leu Gly Gly Leu Ser Leu Gln Glu Val Thr Ser Leu Ala Met
134  440                 445                 450                 455
135  gag gaa tcc caa gaa gca aaa tca ttg cac cag ccc ctg ggg att tgc    1446
136  Glu Glu Ser Gln Glu Ala Lys Ser Leu His Gln Pro Leu Gly Ile Cys
137                  460                 465                 470
138  aca gac aga aca tct gac cca aat gtg cta cac agt ggg gag gaa ggg    1494
139  Thr Asp Arg Thr Ser Asp Pro Asn Val Leu His Ser Gly Glu Glu Gly
140              475                 480                 485
141  aca cca cag tac cta aag ggc cag ctc ccc ctc ctc tcc tca gtc cag    1542
142  Thr Pro Gln Tyr Leu Lys Gly Gln Leu Pro Leu Leu Ser Ser Val Gln
143          490                 495                 500
144  atc gag ggc cac ccc atg tcc ctc cct ttg caa cct cct tcc ggt cca    1590
145  Ile Glu Gly His Pro Met Ser Leu Pro Leu Gln Pro Pro Ser Gly Pro
146      505                 510                 515
147  tgt tcc ccc tcg gat caa ggt cca agt ccc tgg ggc ctg ctg gag tcc    1638
148  Cys Ser Pro Ser Asp Gln Gly Pro Ser Pro Trp Gly Leu Leu Glu Ser
149  520                 525                 530                 535
150  ctt gtg tgt ccc aag gat gaa gcc aag agc cca gcc cct gag acc tca    1686
151  Leu Val Cys Pro Lys Asp Glu Ala Lys Ser Pro Ala Pro Glu Thr Ser
152                  540                 545                 550
153  gac ctg gag cag cct aca gaa ctg gat tct ctt ttc aga ggc ctg gcc    1734
154  Asp Leu Glu Gln Pro Thr Glu Leu Asp Ser Leu Phe Arg Gly Leu Ala
155              555                 560                 565
156  ctg act gtg cag tgg gag tcc tgaggggaat gggaaaggct tggtgcttcc       1785
157  Leu Thr Val Gln Trp Glu Ser
158          570
159  tccctgtccc taccccagtgt cacatccttg gctgtcaatc ccatgcctgc ccatgccaca  1845
160  cactctgcga tctggcctca gacgggtgcc cttgagagaa gcagagggag tggcatgcag  1905
161  ggccctgcgc atgggtgcgc tcttcaccgg aacaaagcag catgataagg actgcagcgg  1965
162  gggagctctg gggagcagct tgtgtagaca agcggtgtgt cgctgagccc tgcaaggcag  2025
163  aaatgacagt gcaaggagga aatgcaggga aactcccgag gtccagagcc ccacctccta  2085
164  acaccatgga ttcaaaagtgc tcagggaatt tgcctctcct tgccccattc ctggccagtt  2145
165  tcacaatcta gctcgacaga gcatgaggcc cctgcctctt ctgtcattgt tcaaaggtgg  2205
166  gaagagagcc tggaaaagaa ccaggcctgg aaaagaacca gaaggaggct gggcagaacc  2265
167  agaacaacct gcacttctgc caaggccagg gccagcagga cggcaggact ctagggaggg  2325
168  gtgtggcctg cagctcattc ccagccaggg caactgcctg acgttgcacg atttcagctt  2385
```

RAW SEQUENCE LISTING DATE: 03/17/2003
PATENT APPLICATION: **US/09/925,055D** TIME: 15:20:10

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\03172003\I925055D.raw**

cattcctctg atagaacaaa gcgaaatcca ggtccaccag ggagggagac acacaagcct 2445
tttctgcagg cagqagtttc agaccctatc ctgagaatgg ggtttgaaag gaaggtgagg 2505
gctgtggccc ctggacgggt acaataacac actgtactga tgtcacaact ttgcaagctc 2565
tgccttgggt tcagcccatc tgggctcaaa ttccagcctc accactcaca agctgtgtga 2625
cttcaaacaa atgaaatcag tgcccagaac ctcggtttcc tcatctgtaa tgtgggatc 2685
ataacaccta cctcatggag ttgtcgtgaa catgaaatga agtcatgtct ttaaagtgct 2745
taatagtgcc tggtacatgg gcagtgccca ataaacggta gctatttaaa aaaaaaaaaa 2805
aaaaaaaaaa atagcggccg cctcga 2831

<210> SEQ ID NO: 2
<211> LENGTH: 574
<212> TYPE: PRT
<213> ORGANISM: Homo sapien
<400> SEQUENCE: 2

Met Arg Thr Leu Leu Thr Ile Leu Thr Val Gly Ser Leu Ala Ala His
1 5 10 15
Ala Pro Glu Asp Pro Ser Asp Leu Leu Gln His Val Lys Phe Gln Ser
20 25 30
Ser Asn Phe Glu Asn Ile Leu Thr Trp Asp Ser Gly Pro Glu Gly Thr
35 40 45
Pro Asp Thr Val Tyr Ser Ile Glu Tyr Lys Thr Tyr Gly Glu Arg Asp
50 55 60
Trp Val Ala Lys Lys Gly Cys Gln Arg Ile Thr Arg Lys Ser Cys Asn
65 70 75 80
Leu Thr Val Glu Thr Gly Asn Leu Thr Glu Leu Tyr Tyr Ala Arg Val
85 90 95
Thr Ala Val Ser Ala Gly Gly Arg Ser Ala Thr Lys Met Thr Asp Arg
100 105 110
Phe Ser Ser Leu Gln His Thr Thr Leu Lys Pro Pro Asp Val Thr Cys
115 120 125
Ile Ser Lys Val Arg Ser Ile Gln Met Ile Val His Pro Thr Pro Thr
130 135 140
Pro Ile Arg Ala Gly Asp Gly His Arg Leu Thr Leu Glu Asp Ile Phe
145 150 155 160
His Asp Leu Phe Tyr His Leu Glu Leu Gln Val Asn Arg Thr Tyr Gln
165 170 175
Met His Leu Gly Gly Lys Gln Arg Glu Tyr Glu Phe Phe Gly Leu Thr
180 185 190
Pro Asp Thr Glu Phe Leu Gly Thr Ile Met Ile Cys Val Pro Thr Trp
195 200 205
Ala Lys Glu Ser Ala Pro Tyr Met Cys Arg Val Lys Thr Leu Pro Asp
210 215 220
Arg Thr Trp Thr Tyr Ser Phe Ser Gly Ala Phe Leu Phe Ser Met Gly
225 230 235 240
Phe Leu Val Ala Val Leu Cys Tyr Leu Ser Tyr Arg Tyr Val Thr Lys
245 250 255
Pro Pro Ala Pro Pro Asn Ser Leu Asn Val Gln Arg Val Leu Thr Phe
260 265 270
Gln Pro Leu Arg Phe Ile Gln Glu His Val Leu Ile Pro Val Phe Asp
275 280 285

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/925,055D

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\03172003\I925055D.raw

```
Leu Ser Gly Ile Ser Ser Leu Ala Gln Pro Val Gln Tyr Ser Gln Ile
      290                 295                 300
Arg Val Ser Gly Pro Arg Glu Pro Ala Gly Ala Pro Gln Arg His Ser
305                 310                 315                 320
Leu Ser Glu Ile Thr Tyr Leu Gly Gln Pro Asp Ile Ser Ile Leu Gln
                325                 330                 335
Pro Ser Asn Val Pro Pro Pro Gln Ile Leu Ser Pro Leu Ser Tyr Ala
            340                 345                 350
Pro Asn Ala Ala Pro Glu Val Gly Pro Pro Ser Tyr Ala Pro Gln Val
        355                 360                 365
Thr Pro Glu Ala Gln Phe Pro Phe Tyr Ala Pro Gln Ala Ile Ser Lys
    370                 375                 380
Val Gln Pro Ser Ser Tyr Ala Pro Gln Ala Thr Pro Asp Ser Trp Pro
385                 390                 395                 400
Pro Ser Tyr Gly Val Cys Met Glu Gly Ser Gly Lys Asp Ser Pro Thr
                405                 410                 415
Gly Thr Leu Ser Ser Pro Lys His Leu Arg Pro Lys Gly Gln Leu Gln
            420                 425                 430
Lys Glu Pro Pro Ala Gly Ser Cys Met Leu Gly Gly Leu Ser Leu Gln
        435                 440                 445
Glu Val Thr Ser Leu Ala Met Glu Glu Ser Gln Glu Ala Lys Ser Leu
    450                 455                 460
His Gln Pro Leu Gly Ile Cys Thr Asp Arg Thr Ser Asp Pro Asn Val
465                 470                 475                 480
Leu His Ser Gly Glu Glu Gly Thr Pro Gln Tyr Leu Lys Gly Gln Leu
                485                 490                 495
Pro Leu Leu Ser Ser Val Gln Ile Glu Gly His Pro Met Ser Leu Pro
            500                 505                 510
Leu Gln Pro Pro Ser Gly Pro Cys Ser Pro Ser Asp Gln Gly Pro Ser
        515                 520                 525
Pro Trp Gly Leu Leu Glu Ser Leu Val Cys Pro Lys Asp Glu Ala Lys
    530                 535                 540
Ser Pro Ala Pro Glu Thr Ser Asp Leu Glu Gln Pro Thr Glu Leu Asp
545                 550                 555                 560
Ser Leu Phe Arg Gly Leu Ala Leu Thr Val Gln Trp Glu Ser
                565                 570

<210> SEQ ID NO: 3
<211> LENGTH: 211
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 3
 Pro Glu Asp Pro Ser Asp Leu Leu Gln His Val Lys Phe Gln Ser Ser
  1               5                  10                  15
Asn Phe Glu Asn Ile Leu Thr Trp Asp Ser Gly Pro Glu Gly Thr Pro
            20                  25                  30
Asp Thr Val Tyr Ser Ile Glu Tyr Lys Thr Tyr Gly Glu Arg Asp Trp
        35                  40                  45
Val Ala Lys Lys Gly Cys Gln Arg Ile Thr Arg Lys Ser Cys Asn Leu
    50                  55                  60
```

VERIFICATION SUMMARY DATE: 03/17/2003
PATENT APPLICATION: US/09/925,055D TIME: 13:20:11

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\03172003\I925055D.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:953 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:956 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29

OIPE

RAW SEQUENCE LISTING DATE: 03/12/2003
PATENT APPLICATION: US/09/925,055D TIME: 08:57:36

Input Set : A:\00-56 SEQ.txt
Output Set: N:\CRF4\03122003\I925055D.raw

4 <110> APPLICANT: Kindsvogel, Wayne R.
5 Topouzis, Stavros
6 <120> TITLE OF INVENTION: SOLUBLE ZCYTOR11 CYTOKINE RECEPTORS
7 <130> FILE REFERENCE: 00-56
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/925,055D
C--> 14 <141> CURRENT FILING DATE: 2003-03-04
12 <150> PRIOR APPLICATION NUMBER: US 60/223,827
13 <151> PRIOR FILING DATE: 2000-08-08
14 <150> PRIOR APPLICATION NUMBER: US 60/250,876
15 <151> PRIOR FILING DATE: 2000-12-01
16 <160> NUMBER OF SEQ ID NOS: 35
17 <170> SOFTWARE: PatentIn for Windows Version 3.0

ERRORED SEQUENCES

Does Not Comply
Corrective Diskette Needed

1.45 <210> SEQ ID NO: 35
1.46 <211> LENGTH: 201
1.47 <212> TYPE: PRT
1.48 <213> ORGANISM: Homo sapiens
1.50 <400> SEQUENCE: 35

```
1.51  Asp Glu Val Ala Ile Leu Pro Ala Pro Gln Asn Leu Ser Val Leu Ser
1.52   1               5                  10                  15
1.53  Thr Asn Met Lys His Leu Leu Met Trp Ser Pro Val Ile Ala Pro Gly
1.54              20                  25                  30
1.55  Glu Thr Val Tyr Tyr Ser Val Glu Tyr Gln Gly Glu Tyr Glu Ser Leu
1.56          35                  40                  45
1.57  Tyr Thr Ser His Ile Trp Ile Pro Ser Ser Trp Cys Ser Leu Thr Glu
1.58      50                  55                  60
1.59  Gly Pro Glu Cys Asp Val Thr Asp Asp Ile Thr Ala Thr Val Pro Tyr
1.60  65                  70                  75                  80
1.61  Asn Leu Arg Val Arg Ala Thr Leu Gly Ser Gln Thr Ser Ala Trp Ser
1.62                  85                  90                  95
1.63  Ile Leu Lys His Pro Phe Asn Arg Asn Ser Thr Ile Leu Thr Arg Pro
1.64              100                 105                 110
1.65  Gly Met Glu Ile Thr Lys Asp Gly Phe His Leu Val Ile Glu Leu Glu
1.66          115                 120                 125
1.67  Asp Leu Gly Pro Gln Phe Glu Phe Leu Val Ala Tyr Trp Arg Arg Glu
1.68      130                 135                 140
1.69  Pro Gly Ala Glu Glu His Val Lys Met Val Arg Ser Gly Gly Ile Pro
1.70  145                 150                 155                 160
1.71  Val His Leu Glu Thr Met Glu Pro Gly Ala Ala Tyr Cys Val Lys Ala
1.72                  165                 170                 175
```

RAW SEQUENCE LISTING DATE: [illegible]
PATENT APPLICATION: US/09/925,055D TIME: [illegible]

Input Set : A:\00-56 SEQ.txt
Output Set: N:\CRF4\03122003\I925055D.raw

```
1273  Gln Thr Phe Val Lys Ala Ile Gly Arg Tyr Ser Ala Pro Ser Gln Thr
1274                  180                 185                 190
1275  Glu Cys Val Glu Val Gln Gly Glu Ala
1276          195                 200
```

E--> 1280 30

delete

VERIFICATION SUMMARY DATE: [illegible]
PATENT APPLICATION: US/09/925,055D TIME: [illegible]

Input Set : A:\00-56 SEQ.txt
Output Set: N:\CRF4\03122003\I925055D.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:956 M:251 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:956 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29
L:1280 M:332 E: (2) Invalid/Missing Amino Acid Numbering, SEQ ID:65